

Table S5. Extremely conserved regions compared to the remaining nucleotides in each of three satDNA families.

Satellite	Region: Start-End	Conservation	Homozygosity	P-value
VspSat02-101	49-59: AHCTTAHCCCC	0,818	0,990	0,0000
VspSat04-107	46-55: ACAAAMATAT	0,900	0,986	0,0001
VspSat05-141	106-124: GATTGATCTAGCACCACTT	1,000	1,000	0,0003